

SEQUENCE LISTING

<110> Ban, Kazuhiro  
Shiotsuka, Hidenori  
Imamura, Takeshi

<120> Kit for immobilizing organic substance, organic substance-immobilized structure, and manufacturing methods therefor

<130> 03500.102556

<140> PCT/JP2005/001316

<141> 2005-01-25

<150> JP2004-016858

<151> 2004-01-26

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 ggacgcgtgt tgctgggga caccctgcat cccaccaacc cgcaagaccg tcgttcgac 240  
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 Ala Ser Ala Arg Met Val Leu Arg Gln Ala Ile Lys Gln Pro Val His  
 35 40 45  
 Ser Val Lys His Val Ala His Phe Gly Leu Glu Leu Lys Asn Val Leu  
 50 55 60  
 Leu Gly Lys Ser Gly Leu Gln Pro Thr Ser Asp Asp Arg Arg Phe Ala  
 65 70 75 80  
 Asp Pro Ala Trp Ser Gln Asn Pro Leu Tyr Lys Arg Tyr Leu Gln Thr  
 85 90 95  
 Tyr Leu Ala Trp Arg Lys Glu Leu His Asp Trp Ile Asp Glu Ser Asn  
 100 105 110  
 Leu Ala Pro Lys Asp Val Ala Arg Gly His Phe Val Ile Asn Leu Met  
 115 120 125  
 Thr Glu Ala Met Ala Pro Thr Asn Thr Ala Ala Asn Pro Ala Ala Val  
 130 135 140

Lys Arg Phe Phe Glu Thr Gly Gly Lys Ser Leu Leu Asp Gly Leu Ser  
 145 150 155 160  
 His Leu Ala Lys Asp Leu Val His Asn Gly Gly Met Pro Ser Gln Val  
 165 170 175  
 Asn Met Gly Ala Phe Glu Val Gly Lys Ser Leu Gly Val Thr Glu Gly  
 180 185 190  
 Ala Val Val Phe Arg Asn Asp Val Leu Glu Leu Ile Gln Tyr Lys Pro  
 195 200 205  
 Thr Thr Glu Gln Val Tyr Glu Arg Pro Leu Leu Val Val Pro Pro Gln  
 210 215 220  
 Ile Asn Lys Phe Tyr Val Phe Asp Leu Ser Pro Asp Lys Ser Leu Ala  
 225 230 235 240  
 Arg Phe Cys Leu Arg Asn Asn Val Gln Thr Phe Ile Val Ser Trp Arg  
 245 250 255  
 Asn Pro Thr Lys Glu Gln Arg Glu Trp Gly Leu Ser Thr Tyr Ile Glu  
 260 265 270  
 Ala Leu Lys Glu Ala Val Asp Val Val Thr Ala Ile Thr Gly Ser Lys  
 275 280 285  
 Asp Val Asn Met Leu Gly Ala Cys Ser Gly Gly Ile Thr Cys Thr Ala  
 290 295 300  
 Leu Leu Gly His Tyr Ala Ala Ile Gly Glu Asn Lys Val Asn Ala Leu  
 305 310 315 320  
 Thr Leu Leu Val Ser Val Leu Asp Thr Thr Leu Asp Ser Asp Val Ala  
 325 330 335  
 Leu Phe Val Asn Glu Gln Thr Leu Glu Ala Ala Lys Arg His Ser Tyr  
 340 345 350  
 Gln Ala Gly Val Leu Glu Gly Arg Asp Met Ala Lys Val Phe Ala Trp  
 355 360 365  
 Met Arg Pro Asn Asp Leu Ile Trp Asn Tyr Trp Val Asn Asn Tyr Leu  
 370 375 380  
 Leu Gly Asn Glu Pro Pro Val Phe Asp Ile Leu Phe Trp Asn Asn Asp  
 385 390 395 400  
 Thr Thr Arg Leu Pro Ala Ala Phe His Gly Asp Leu Ile Glu Leu Phe  
 405 410 415  
 Lys Asn Asn Pro Leu Ile Arg Pro Asn Ala Leu Glu Val Cys Gly Thr  
 420 425 430  
 Pro Ile Asp Leu Lys Gln Val Thr Ala Asp Ile Phe Ser Leu Ala Gly  
 435 440 445  
 Thr Asn Asp His Ile Thr Pro Trp Lys Ser Cys Tyr Lys Ser Ala Gln  
 450 455 460  
 Leu Phe Gly Gly Asn Val Glu Phe Val Leu Ser Ser Ser Gly His Ile  
 465 470 475 480

Gln Ser Ile Leu Asn Pro Pro Gly Asn Pro Lys Ser Arg Tyr Met Thr  
485 490 495

Ser Thr Glu Val Ala Glu Asn Ala Asp Glu Trp Gln Ala Asn Ala Thr  
500 505 510

Lys His Thr Asp Ser Trp Trp Leu His Trp Gln Ala Trp Gln Ala Gln  
515 520 525

Arg Ser Gly Glu Leu Lys Lys Ser Pro Thr Lys Leu Gly Ser Lys Ala  
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<213> Pseudomonas cichorii YN2 ; FERM BP-7375

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Met Arg Asp Lys Pro Ala Arg Glu Ser Leu Pro Thr Pro Ala Lys Phe  
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Ser Thr Leu Arg Ser Val Ala Ala His Gly Leu Arg His Pro Val His  
35 40 45

Thr Ala Arg His Ala Leu Lys Leu Gly Gly Gln Leu Gly Arg Val Leu  
50 55 60

Leu Gly Asp Thr Leu His Pro Thr Asn Pro Gln Asp Arg Arg Phe Asp  
65 70 75 80

Asp Pro Ala Trp Ser Leu Asn Pro Phe Tyr Arg Arg Ser Leu Gln Ala  
85 90 95

Tyr Leu Ser Trp Gln Lys Gln Val Lys Ser Trp Ile Asp Glu Ser Asn  
100 105 110

Met Ser Pro Asp Asp Arg Ala Arg Ala His Phe Ala Phe Ala Leu Leu  
115 120 125

Asn Asp Ala Val Ser Pro Ser Asn Ser Leu Leu Asn Pro Leu Ala Ile  
130 135 140

Lys Glu Ile Phe Asn Ser Gly Gly Asn Ser Leu Val Arg Gly Ile Gly  
145 150 155 160

His Leu Val Asp Asp Leu Leu His Asn Asp Gly Leu Pro Arg Gln Val  
165 170 175

Thr Arg His Ala Phe Glu Val Gly Lys Thr Val Ala Thr Thr Thr Gly  
180 185 190

Ala Val Val Phe Arg Asn Glu Leu Leu Glu Leu Ile Gln Tyr Lys Pro  
195 200 205

Met Ser Glu Lys Gln Tyr Ser Lys Pro Leu Leu Val Val Pro Pro Gln  
210 215 220

Ile Asn Lys Tyr Tyr Ile Phe Asp Leu Ser Pro His Asn Ser Phe Val			
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Gln Phe Ala Leu Lys Asn Gly Leu Gln Thr Phe Val Ile Ser Trp Arg			
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Asn Pro Asp Val Arg His Arg Glu Trp Gly Leu Ser Thr Tyr Val Glu			
	260	265	270
Ala Val Glu Glu Ala Met Asn Val Cys Arg Ala Ile Thr Gly Ala Arg			
	275	280	285
Glu Val Asn Leu Met Gly Ala Cys Ala Gly Gly Leu Thr Ile Ala Ala			
	290	295	300
Leu Gln Gly His Leu Gln Ala Lys Arg Gln Leu Arg Arg Val Ser Ser			
305	310	315	320
Ala Thr Tyr Leu Val Ser Leu Leu Asp Ser Gln Leu Asp Ser Pro Ala			
	325	330	335
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Tyr Gln Lys Gly Val Leu Glu Gly Arg Asp Met Ala Lys Val Phe Ala			
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Trp Met Arg Pro Asn Asp Leu Ile Trp Ser Tyr Phe Val Asn Asn Tyr			
	370	375	380
Leu Met Gly Lys Glu Pro Pro Ala Phe Asp Ile Leu Tyr Trp Asn Asn			
385	390	395	400
Asp Asn Thr Arg Leu Pro Ala Ala Leu His Gly Asp Leu Leu Asp Phe			
	405	410	415
Phe Lys His Asn Pro Leu Ser His Pro Gly Gly Leu Glu Val Cys Gly			
	420	425	430
Thr Pro Ile Asp Leu Gln Lys Val Thr Val Asp Ser Phe Ser Val Ala			
	435	440	445
Gly Ile Asn Asp His Ile Thr Pro Trp Asp Ala Val Tyr Arg Ser Thr			
	450	455	460
Leu Leu Leu Gly Gly Glu Arg Arg Phe Val Leu Ala Asn Ser Gly His			
465	470	475	480
Val Gln Ser Ile Leu Asn Pro Pro Asn Asn Pro Lys Ala Asn Tyr Leu			
	485	490	495
Glu Gly Ala Lys Leu Ser Ser Asp Pro Arg Ala Trp Tyr Tyr Asp Ala			
	500	505	510
Lys Pro Val Asp Gly Ser Trp Trp Thr Gln Trp Leu Gly Trp Ile Gln			
	515	520	525
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 <223> Complimentary chain for ssDNA of SEQ ID:3  
  
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 <400> 53  
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<210> 54  
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<223> Coding chain for peptide of SEQ ID:5

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<210> 56  
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<223> Complimentary chain for ssDNA of SEQ ID:5

<400> 56  
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<400> 57  
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<210> 58  
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<400> 58  
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<210> 59  
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<223> Coding chain for peptide of SEQ ID:7



<400> 59  
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<210> 60  
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<210> 61  
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<223> Coding chain for peptide of SEQ ID:8

<400> 61  
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<210> 62  
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<210> 63  
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<210> 64  
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<400> 64  
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<210> 65  
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<400> 65

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<210> 66

<211> 50

<212> DNA

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<223> Complimentary chain for ssDNA of SEQ ID:10

<400> 66

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<210> 67

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:11

<400> 67

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<210> 68

<211> 50

<212> DNA

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<223> Complimentary chain for ssDNA of SEQ ID:11

<400> 68

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<210> 69

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:12

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<210> 70

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<212> DNA

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<223> Complimentary chain for ssDNA of SEQ ID:12

<400> 70

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<210> 71

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 <400> 71  
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 <210> 72  
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 <223> Complimentary chain for ssDNA of SEQ ID:13  
  
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 <210> 73  
 <211> 58  
 <212> DNA  
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 <220>  
 <223> Coding chain for peptide of SEQ ID:14  
  
 <400> 73  
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 <210> 74  
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 <223> Complimentary chain for ssDNA of SEQ ID:14  
  
 <400> 74  
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 <210> 75  
 <211> 58  
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 <223> Coding chain for peptide of SEQ ID:15  
  
 <400> 75  
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 <210> 76  
 <211> 50  
 <212> DNA  
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 <223> Complimentary chain for ssDNA of SEQ ID:15  
  
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<210> 77

<211> 58

<212> DNA

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<223> Coding chain for peptide of SEQ ID:16

<400> 77

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<210> 78

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Complimentary chain for ssDNA of SEQ ID:16

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<211> 58

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<223> Coding chain for peptide of SEQ ID:17

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<210> 80

<211> 50

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<223> Complimentary chain for ssDNA of SEQ ID:17

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<210> 81

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<210> 82

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<223> Complimentary chain for ssDNA of SEQ ID:18

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<210> 84

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<223> Complimentary chain for ssDNA of SEQ ID:19

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<210> 85

<211> 58

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<223> Coding chain for peptide of SEQ ID:20

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<210> 86

<211> 50

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<223> Complimentary chain for ssDNA of SEQ ID:20

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<210> 87

<211> 58

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<223> Coding chain for peptide of SEQ ID:21

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<210> 88

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<223> Complimentary chain for ssDNA of SEQ ID:21

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<211> 58

<212> DNA

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<223> Coding chain for peptide of SEQ ID:22

<400> 89

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<210> 90

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Complimentary chain for ssDNA of SEQ ID:22

<400> 90

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<210> 91

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:23

<400> 91

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<210> 92

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Complimentary chain for ssDNA of SEQ ID:23

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<210> 93

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:24

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<210> 94  
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<400> 94  
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<210> 95  
<211> 58  
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<210> 96  
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<210> 98  
<211> 50  
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<210> 100  
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<220>  
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<210> 102  
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<400> 102  
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<210> 103  
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<210> 104  
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<220>  
<223> Coding chain for peptide of SEQ ID:30  
  
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<212> DNA  
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<223> Complimentary chain for ssDNA of SEQ ID:30  
  
<400> 106  
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<210> 107  
<211> 52  
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<220>  
<223> Coding chain for peptide of SEQ ID:31  
  
<400> 107  
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<210> 108  
<211> 44  
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<223> Complimentary chain for ssDNA of SEQ ID:31  
  
<400> 108  
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<210> 109  
<211> 52  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<223> Coding chain for peptide of SEQ ID:32  
  
<400> 109  
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<210> 110  
<211> 44  
<212> DNA  
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<210> 111  
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<220>  
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gagctaagat cagaccctcg tattgccgag agcatccttc gtcttcactt ccacgactgc 180  
ttgttaatg gttgtgacgc atcgatcttg ttagacaaca caacatcatt tgaacagag 240  
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gccgcggtgg agagtgcag cccaagaacc gtttcatgag cagatttgc caccattgca 360  
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gcactgtccg ggggccacac atttggtaaa aatcagtgtc ggtttattat ggacagatta 600  
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aactccaact ct 972

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<220>  
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<210> 114

<211> 120

<212> DNA

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<223> Primer for PCR multiplication

<400> 114

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<212> DNA

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<223> Primer for PCR multiplication

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<223> Primer for PCR multiplication

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<223> Primer for PCR multiplication

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<210> 124  
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<223> Primer for PCR multiplication

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<210> 125

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<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 125

tacaactca gcaacaccgg ttaccgat 30

<210> 126

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<223> Primer for PCR multiplication

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<223> Primer for PCR multiplication

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<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

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agagttggag ttcaccaccc tacaattcaa 30

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<212> DNA

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agtcggatcc gtttatgcga atcagactcc gccttctaag gcgcgggggtg gaggttcg 58

<210> 133

<211> 34

<212> DNA

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<223> Primer for PCR multiplication

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<212> DNA

<213> Artificial Sequence

<220>

<223> GroEL coding artificial sense-sequence

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gacgtaaaat tcggtaacga cgctcgtgtg aaaatgctgc gcggcgtaaa cgtactggca 120

gatgcagtga aagtaccct cgttccaaaa ggccgtaacg tagttctgga taaatcttc 180  
 ggtgcaccga ccatcaccaa agatggtgtt tccgttgctc gtgaaatcga actggaagac 240  
 aagttcgaat atatgggtgc gcagatggtg aaagaagttg cctctaaagc aaacgacgct 300  
 gcaggcgacg gtaccaccac tgcaaccgta ctggctcagg ctatcatcac tgaaggtctg 360  
 aaagctgttg ctgcgggcat gaacccgatg gacctgaaac gtggtatcga caaagcggtt 420  
 accgtgcag ttgaagaact gaaagcgtg tccgtacat gctctgactc taaagcgatt 480  
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 gacgaactgg acgtggttga aggtatgcag ttcgaccgtg gctacctgc tcctacttc 660  
 atcaacaagc cggaaactgg cgcagtagaa ctggaagcc cgttcacct gctggctgac 720  
 aagaaaatct ccaacatccg cgaatgctg ccggttctg aagctgttc caaagcaggc 780  
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 cgtaaagcta tgctgcagga tatcgcaacc ctgactggcg gtaccgtgat ctctgaagag 960  
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 cgtgttgctc agatccgtca gcagattgaa gaagcaact ctgactacga ccgtgaaaa 1140  
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 accgaagtgt aatgaaaga gaaaaagca cgcgttgaag atgccctgca cgcgacctgt 1260  
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 tacggcaaca tgatgacat gggatcctg gaccaacca aagtaactc ttctgctctg 1560  
 cagtacgcag cttctgtggc tggcctgat atcaccaccg aatgcatggt taccgacctg 1620  
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<211> 120

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

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<210> 136  
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<220>  
<223> Primer for PCR multiplication

<400> 136  
gtttatgcga atcagactcc gccttctaag 30

<210> 137  
<211> 120  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

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cgttacggcc ttttgaccg agggttaactt tcaactgcac tgccagtacg ttacgccgc 120

<210> 138  
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<220>  
<223> Primer for PCR multiplication

<400> 138  
gagcaacgga aacaccatct ttggtgatgg 30

<210> 139  
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<220>  
<223> Primer for PCR multiplication

<400> 139  
agatggtgtt tccgttgctc gtgaaatcga actggaagac aagttcgaaa atatgggtgc 60  
gcagatggtg aaagaagttg cctctaaagc aaacgacgct gcaggcgacg gtaccaccac 120

<210> 140  
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<220>  
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<400> 140  
agatggtgtt tccgttgctc gtgaaatcga 30



<210> 141  
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<220>  
<223> Primer for PCR multiplication

<400> 141  
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<210> 142  
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<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 142  
aaccgctttg tcgataccac gtttcaggtc 30

<210> 143  
<211> 120  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 143  
gtggtatcga caaagcggt accgctgcag ttgaagaact gaaagcgctg tccgtacat 60  
gctctgactc taaagcgatt gctcaggtg gtaccatctc cgctaactcc gacgaaaccg 120

<210> 144  
<211> 30  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 144  
gtggtatcga caaagcggt accgctgcag 30

<210> 145  
<211> 120  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 145  
tcaaccagt ccagttcgtc ctgcagaccg gtaccgtctt caacgggat aacgccttct 60  
ttaccgactt tgccatcgc ttcagcagtc agttaccta cggtttcgtc ggagttagcg 120

<210> 146

<211> 30  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 146  
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<210> 147  
<211> 120  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 147  
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atcaacaagc cggaaactgg cgcagtagaa ctggaaagcc cgttcacatc gctggctgac 120

<210> 148  
<211> 30  
<212> DNA  
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<220>  
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<400> 148  
gacgaactgg acgtggttga aggtatgcag 30

<210> 149  
<211> 120  
<212> DNA  
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<220>  
<223> Primer for PCR multiplication

<400> 149  
cttcgccttc tacatcttca gcgatgataa gcagcggttt gcctgctttg gcaacagctt 60  
ccagaaccgg cagcatttcg cggatgttgg agattttctt gtcagccagc aggatgaacg 120

<210> 150  
<211> 30  
<212> DNA  
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<220>  
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<400> 150  
cttcgccttc tacatcttca gcgatgataa 30

<210> 151  
<211> 120  
<212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 151  
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<210> 152  
 <211> 30  
 <212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 152  
 tgaagatgta gaaggcgaag cgctggcaac 30

<210> 153  
 <211> 120  
 <212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 153  
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 ctcttcagag atcacggtag cgccagtcag gggtgcgata tcttcagca tagctttacg 120

<210> 154  
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<220>  
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<400> 154  
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<210> 155  
 <211> 120  
 <212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 155  
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 aagaagctgc aatccagggc cgtgttgctc agatccgtca gcagattgaa gaagcaactt 120

<210> 156  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer for PCR multiplication

<400> 156  
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<210> 157  
 <211> 120  
 <212> DNA  
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<220>  
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<400> 157  
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<210> 158  
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 <212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 158  
 tcttcattt caactcggg agcagcacc 30

<210> 159  
 <211> 120  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer for PCR multiplication

<400> 159  
 accgaagtg aatgaaaga gaaaaagca cgcgtgaag atgccctgca cgcgaccgt 60  
 gctgcgtag aagaaggcgt ggtgctggt ggtggtgtg cgtgatccg cgtacgtct 120

<210> 160  
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 <212> DNA  
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<220>  
 <223> Primer for PCR multiplication

<400> 160  
 accgaagtg aatgaaaga gaaaaagca 30

<210> 161  
 <211> 120  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer for PCR multiplication

<400> 161  
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cggtctggtc ttggttctga ccacgcaggt cagccagttt agacgctacg cggatcagcg 120

<210> 162

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 162

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<210> 163

<211> 120

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 163

gcgtcagatc gtattgaact gcggcgaaga accgtctgtt gttgctaaca ccgttaaagg 60

cggcgacggc aactacgggtt acaacgcagc aaccgaagaa tacggcaaca tgatcgacat 120

<210> 164

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 164

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<210> 165

<211> 120

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 165

caggtcggta accatgcatt cggtggtgat catcaggcca gccacagaag ctgcgtactg 60

cagagcagaa cgagttactt tggttgggtc caggataccc atgtcgatca tgttgccgta 120

<210> 166

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer for PCR multiplication

<400> 166

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<210> 167

<211> 95

<212> DNA  
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 <223> Primer for PCR multiplication  
  
 <400> 167  
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 tgcacgttt ttcggcaggt cggtaacct gcatt 95  
  
 <210> 168  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence  
  
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 <223> Primer for PCR multiplication  
  
 <400> 168  
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 <210> 169  
 <211> 33  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Primer for PCR multiplication  
  
 <400> 169  
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 <210> 170  
 <211> 8  
 <212> PRT  
 <213> Artificial Sequence  
  
 <220>  
 <223> anodisk membrane-binding peptide  
  
 <400> 170  
 Tyr Ala Gln Thr Pro Pro Ser Arg  
 1 5  
  
 <210> 171  
 <211> 12  
 <212> PRT  
 <213> Artificial Sequence  
  
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 <400> 171  
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 1 5 10  
  
 <210> 172  
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 <212> PRT  
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<223> anodisk membrane-binding peptide

<400> 172

Val Tyr Ala Asn Gln Thr Pro Pro Ser Arg Ala Arg Ala Lys Ala Arg  
1 5 10 15

<210> 173

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> anodisk membrane-binding peptide

<400> 173

Val Tyr Ala Asn Gln Thr Pro Pro Ser Lys Ala Arg Tyr Ala Gln  
1 5 10 15  
Thr Pro Pro Ser Arg  
20

<210> 174

<211> 46

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:170

<400> 174

gatcctatgc gcagactccg cttctcggg gtggagggtc ggagct 46

<210> 175

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Complimentary chain for ssDNA of SEQ ID:170

<400> 175

ccgaacctcc accccgagaa ggcggagtct gcgcatag 38

<210> 176

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Coding chain for peptide of SEQ ID:171

<400> 176

gatccctcta tgcgcaacag actccgcctt ctcggtctcg gggaggaggt tcggagct 58

<210> 177

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Complimentary chain for ssDNA of SEQ ID:171

<400> 177

ccgaacctcc accccgagac cgagaaggcg gagtctgttg cgcataagag 50

<210> 178  
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<220>  
<223> Coding chain for peptide of SEQ ID:1

<400> 178  
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gttcggagct 70

<210> 179  
<211> 62  
<212> DNA  
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<220>  
<223> Complimentary chain for ssDNA of SEQ ID:1

<400> 179  
ccgaacctcc accccgcgcc ttgcgcgtg cgcgagaagg cggagtctga ttcgcataaa 60  
cg 62

<210> 180  
<211> 82  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Coding chain for peptide of SEQ ID:1

<400> 180  
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ctcggggtgg aggttcggag ct 82

<210> 181  
<211> 74  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Complimentary chain for ssDNA of SEQ ID:1

<400> 181  
ccgaacctcc accccgagaa ggccggagtct gcgcataccg cgccttagaa ggccggagtct 60  
gattcgcata aacg 74